

=> d his

(FILE 'HOME' ENTERED AT 11:08:35 ON 20 MAR 2005)

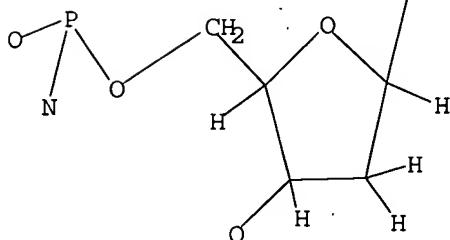
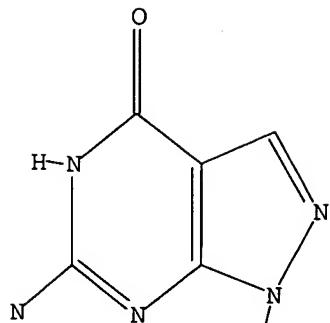
FILE 'REGISTRY' ENTERED AT 11:08:51 ON 20 MAR 2005

L1 STRUCTURE UPLOADED
L2 0 S L1 SSS SAM
L3 0 S L1 SSS FULL
L4 STRUCTURE UPLOADED
L5 6 S L4 SSS SAM
L6 77 S L4 SSS FULL
L7 STRUCTURE UPLOADED
L8 1 S L7 SSS SAM
L9 10 S L7 SSS FULL
L10 STRUCTURE UPLOADED
L11 1 S L10 SSS SAM
L12 9 S L10 SSS FULL
L13 STRUCTURE UPLOADED
L14 1 S L13 SSS SAM
L15 10 S L13 SSS FULL
L16 STRUCTURE UPLOADED
L17 0 S L16 SSS SAM
L18 0 S L16 SSS FULL

FILE 'CAPLUS, MEDLINE' ENTERED AT 11:43:26 ON 20 MAR 2005

L19 4 S L9 AND PHOSPHORAMIDITE?
L20 4 S L12 AND PHOSPHORAMIDITE?
L21 4 S L15 AND PHOSPHORAMIDITE?
L22 0 S L19 AND PHOTOLAB?
L23 0 S L20 AND PHOTOLAB?
L24 0 S L21 AND PHOTOLAB?

=> d 11
L1 HAS NO ANSWERS
L1 STR



Structure attributes must be viewed using STN Express query preparation.

=> s 11 sss sam
SAMPLE SEARCH INITIATED 11:09:29 FILE 'REGISTRY'
SAMPLE SCREEN SEARCH COMPLETED - 7 TO ITERATE

100.0% PROCESSED 7 ITERATIONS 0 ANSWERS
SEARCH TIME: 00.00.01

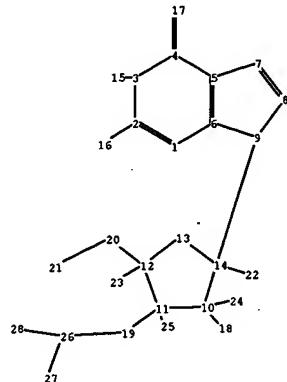
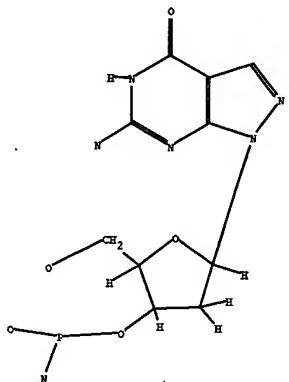
FULL FILE PROJECTIONS: ONLINE **COMPLETE**
BATCH **COMPLETE**
PROJECTED ITERATIONS: 7 TO 298
PROJECTED ANSWERS: 0 TO 0

L2 0 SEA SSS SAM L1

=> s 11 sss full
FULL SEARCH INITIATED 11:09:36 FILE 'REGISTRY'
FULL SCREEN SEARCH COMPLETED - 75 TO ITERATE

100.0% PROCESSED 75 ITERATIONS 0 ANSWERS
SEARCH TIME: 00.00.01

L3 0 SEA SSS FUL L1



chain nodes :

15 16 17 18 19 20 21 22 23 24 25 26 27 28

ring nodes :

1 2 3 4 5 6 7 8 9 10 11 12 13 14
26-27 26-28

chain bonds :

2-16 3-15 4-17 9-14 10-18 10-24 11-19 11-25 12-20 12-23 14-22 19-26 20-21
26-27 26-28

ring bonds :

1-2 1-6 2-3 3-4 4-5 5-6 5-7 6-9 7-8 8-9 10-11 10-14 11-12 12-13 13-14

exact/norm bonds :

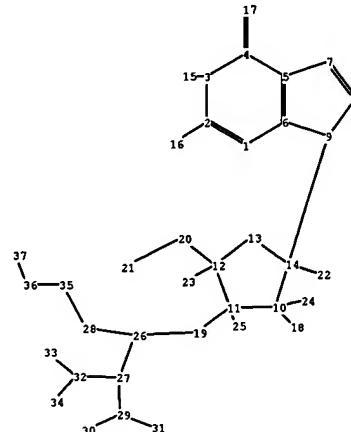
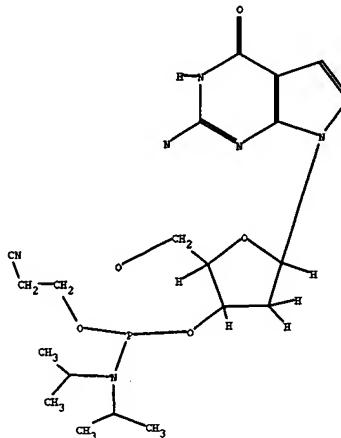
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11-19 12-13 13-14 19-26 26-27 26-28

exact bonds :

3-15 10-18 10-24 11-25 12-20 12-23 14-22 20-21

Match level :

1:Atom 2:Atom 3:Atom 4:Atom 5:Atom 6:Atom 7:Atom 8:Atom 9:Atom 10:Atom 11:Atom
12:Atom 13:Atom 14:Atom 15:CLASS 16:CLASS 17:CLASS 18:CLASS 19:CLASS 20:CLASS
21:CLASS 22:CLASS 23:CLASS 24:CLASS 25:CLASS 26:CLASS 27:CLASS 28:CLASS



chain nodes :

15 16 17 18 19 20 21 22 23 24 25 26 27 28 29 30 31 32 33 34 35 36
37

ring nodes :

1 2 3 4 5 6 7 8 9 10 11 12 13 14

chain bonds :

2-16 3-15 4-17 9-14 10-18 10-24 11-19 11-25 12-20 12-23 14-22 19-26 20-21
26-27 26-28 27-29 27-32 28-35 29-30 29-31 32-33 32-34 35-36 36-37

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exact/norm bonds :

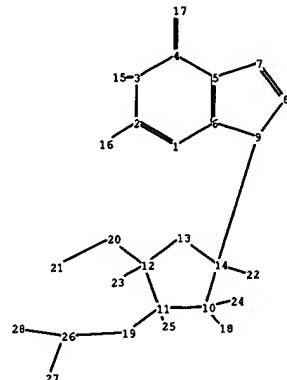
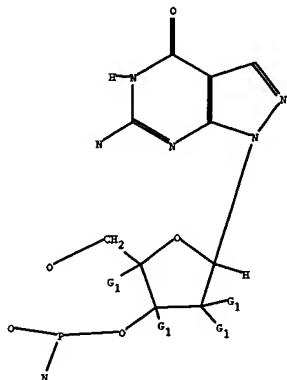
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11-19 12-13 13-14 19-26 26-27 26-28 27-29 27-32

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35-36 36-37

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30:CLASS 31:CLASS 32:CLASS 33:CLASS 34:CLASS 35:CLASS 36:CLASS 37:CLASS



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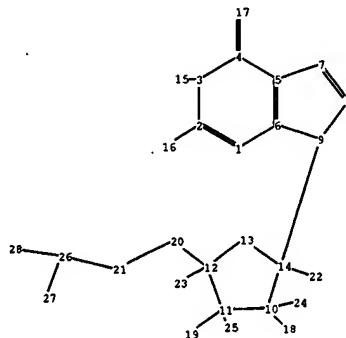
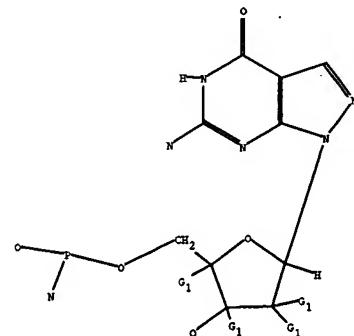
exact bonds :

3-15 12-20 14-22 20-21

G1:H,Cl,Br,F,I,O,Ak

Match level :

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ring nodes :

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